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UPDATE: SECOND ALBANY CONFERENCE ON COMPUTATIONAL BIOLOGY

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The program for the Second Albany Conference on Computational Biology to be held October 8-11, 1992 has been set. The theme of the Conference is "Patterns of Biological Organization". Five sessions are planned that will cover biological self-organization from the level of sequence to the level of embryo development. In addition, workshops are being planned on such topics as data bases and visualization tools. Leading specialists in the various disciplines are being invited, with the degree of involvement in novel computational approaches as one of the most important criteria for selection.

We are seeking an interdisciplinary audience, mathematicians, and computer scientists as well as biologists. All participants will be invited to submit abstracts for posters, although submission of abstracts will not be mandatory.

Sessions will have four speakers each. Topics are:

Session 1 Sequence analysis and secondary structure

Session 2 Tertiary structure prediction

Session 3 Macromolecular function

Session 4 Recognition and assembly

Session 5 Development

Speakers will include:

James Bowie (UCLA)
Steve Bryant (NCBI, NIH)
Jean Michel Claverie (NCBI, NIH)
David DeRossier (Brandeis Univ.)
Rick Fine (Columbia Univ.)
Judith Hempel (BIOSYM)
Michael Levine (UC San Diego)
John Reinitz (Yale Univ.)
Phoebe Stewart (Wistar Inst., Univ. Penn.)
John Sedat (UC San Francisco)
David Waltz (Thinking Machines and Brandeis Univ.)

The registration fee, which includes room and board, is $500. If funding permits, we will sponsor several "young investigator" travel awards to help graduate students and postdocs from the different disciplines attend the conference.

Source: https://www.industrydocuments.ucsf.edu/docs/gxlv0000